

Errata list for paper I

Functional analysis and localisation of a delta-class glutathione S-transferase from *Sarcoptes scabiei*

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Page, line or Table: column	Reads	Should read
41, 50: 1	...the +2 codon was changed to an AAA-codon(lysine)...	... an additional AAA-codon (lysine) was inserted directly after the start codon...
46, Table 2: ssGST3	405 ± 12	4.1 ± 0.1
	193 ± 5.7	1.9 ± 0.1
	689 ± 67	6.9 ± 0.7
46, Table 2: ssGST5	958 ± 39	9.6 ± 0.4
	470 ± 19	4.7 ± 0.2
	1516 ± 196	15.2 ± 2.0
47, 48: 1	...ssGST3 and ssGST5 had relatively high affinity towards CDNB...	...ssGST3 and ssGST5 had relatively low affinity towards CDNB...
47, 1: 2	...ssGST5 had the highest turnover rate ($k_{\text{cat}}=470\pm19\text{ s}^{-1}$)...	...ssGST5 had the highest turnover rate ($k_{\text{cat}}=4.7\pm0.2\text{ s}^{-1}$)...
47, 2: 2	...the most reactive enzyme in catalyzing CDNB conjugation ($k_{\text{cat}}/K_{\text{M}}=1516\pm196\text{ mM}^{-1}\text{ s}^{-1}$).	...the most reactive enzyme in catalyzing CDNB conjugation ($k_{\text{cat}}/K_{\text{M}}=15.2\pm2.0\text{ mM}^{-1}\text{ s}^{-1}$).